

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/019,1148
Source:	PITIO
Date Processed by STIC:	7/30/04

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

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- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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 U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two. Lobby Room 1B03, Arlington, VA 22202

Revised 05/17/04



PCT

RAW SEQUENCE LISTING DATE: 07/30/2004
PATENT APPLICATION: US/10/019,114B TIME: 09:24:30

Input Set : A:\01997.013000 sequence listing.txt

Output Set: N:\CRF4\07302004\J019114B.raw

5 <110> APPLICANT: Seishi KATO

7 Tomoko KIMURA

11 <120> TITLE OF INVENTION: Human proteins having hydrophobic domains and DNAs encoding these

13 proteins

16 <130> FILE REFERENCE: 1997.13000

19 <140> CURRENT APPLICATION NUMBER: US 10/019,114B

C--> 21 <141> CURRENT FILING DATE: 2002-10-24

24 <150> PRIOR APPLICATION NUMBER: JP 11-188835

26 <151> PRIOR FILING DATE: 1999-07-02

30 <160> NUMBER OF SEQ ID NOS: 30

ERRORED SEQUENCES

1769 <210> SEQ ID NO: 24 1771 <211> LENGTH: 1260 1773 <212> TYPE: DNA 1775 <213> ORGANISM: Homo sapiens 1779 <220> FEATURE: 1781 <221> NAME/KEY: CDS 1783 <222> LOCATION: (147)(863) 1787 <400> SEQUENCE: 24												
1789 agetteecee aageggegge ageaceacea geggeageeg eeggageege egeegeageg 60												
1791 gggacgggga gccccggggg gccccgccac tgccgccgtc cgccgtcacc tacccggact 120												
1793 ggatcggcca gagttactcc gaggtg atg agc ctc aac gag cac tcc atg	170											
1795 Met Ser Leu Asn Glu His Ser Met												
1797 1 5												
1799 cag gcg ctg tcc tgg cgc aag ctc tac ttg agc cgc gcc aag ctt aaa	218											
1801 Gln Ala Leu Ser Trp Arg Lys Leu Tyr Leu Ser Arg Ala Lys Leu Lys												
1803 10 15 20												
1805 gcc tcc agc cgg acc tcg gct ctg ctc tcc ggc ttc gcc atg gtg gca	266											
1807 Ala Ser Ser Arg Thr Ser Ala Leu Leu Ser Gly Phe Ala Met Val Ala												
1809 25 30 35 40	314											
1811 atg gtg gag gtg cag ctg gac gct gac cac gac tac cca ccg ggg ctg	314											
1813 Met Val Glu Val Gln Leu Asp Ala Asp His Asp Tyr Pro Pro Gly Leu 1815 45 50 55												
1817 ctc atc gcc ttc agt gcc tgc acc aca gtg ctg gtg gct gtg cac ctg	362											
1819 Leu Ile Ala Phe Ser Ala Cys Thr Thr Val Leu Val Ala Val His Leu	302											
1821 60 65 70												
1823 ttt gcg ctc atg atc agc acc tgc atc ctg ccc aac atc gag gcg gtg	410											
1825 Phe Ala Leu Met Ile Ser Thr Cys Ile Leu Pro Asn Ile Glu Ala Val												
1827 75 80 85												
1829 agc aac gtg cac aat ctc aac tcg gtc aag gag tcc ccc cat gag cgc	458											

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	1831	Ser	Asn	Val	His	Asn	Leu	Asn	Ser	Val	Lys	Glu	Ser	Pro	His	Glu	Arg		
	1833		90					95					100						
	1835	atg	cac	cgc	cac	atc	gag	ctg	gcc	tgg	gcc	ttc	tcc	acc	gtc	atc	ggc	506	
	1837	Met	His	Arg	His	Ile	Glu	Leu	Ala	Trp	Ala	Phe	Ser	Thr	Val	Ile	Gly		
	1839	105					110					115					120		
	1841	acg	ctg	ctc	ttc	cta	gct	gag	gtg	gtg	ctg	ctc	tgc	tgg	gtc	aag	ttc	554	
	1843	Thr	Leu	Leu	Phe	Leu	Àla	Glu	Val	Val	Leu	Leu	Cys	Trp	Val	Lys	Phe	•	
	1845					125					130					135			
	1847	ttg	CCC	ctc	aag	aag	cag	cca	ggc	cag	cca	agg	CCC	acc	agc	aag	CCC	602	
	1849	Leu	Pro	Leu	Lys	Lys	Gln	Pro	Gly	Gln	Pro	Arg	Pro	Thr	Ser	Lys	Pro		
	1851				140					145					150				•
	1853	CCC	gcc	agt	ggc	gca	gca	gcc	aac	gtc	agc	acc	agc	ggc	atc	acc	ccg	650	
	1855	Pro	Ala	Ser	Gly	Ala	Ala	Ala	Asn	Val	Ser	Thr	Ser	Gly	Ile	Thr	Pro		` 1.1
	1857			155	A-				160					165					nooled
E>	1859																	698	huster
	1861	Gly	Gln	Ala	AYa	Ala	Ile	Ala	Ser	Thr	Thr	Ile	Met	Val	Pro	Phe	Gly		nucu
	1863		170					175					180						and.
	1865	_				_		_	_				_		_	_	_	746	mileie aid designation
	1867		Ile	Phe	Ile	Val		Ala	Val	His	Phe	_	Arg	Ser	Leu	Val			any
	1869						190					195					200		
	1871		_		-	_	_		_					_				794	
	1873	His	Lys	Thr	Asp	_	Gln	Phe	Gln	Glu		Asn	Glu	Leu	Ala		Phe		
	1875					205					210					215			
	1877	_	_		_	_	_	_	_		_		-			_	-	842	
	1879	Ala	Arg	Leu		Asp	Gln	Leu	Asp		Arg	Gly	Asp	His		Leu	Thr		
	1881				220					225					230				
	1883								gccca	atg 1	tggt	ctgg	ge e	cttc	cagt	3		890	
	1885	Pro	GLY		His	Tyr	Ala												
	1887			235															
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					_	_	_	_	_		_			_	_	_	ctcaac	1190	
				_	ccag	gcatt	g ac	grgr	gtaca	a tg	gtgt	gtg	acad	cata	aat a	atact	cataa	1250	
	1901	gga	cacct	CC														1260	

VERIFICATION SUMMARY

DATE: 07/30/2004 TIME: 09:24:31

PATENT APPLICATION: US/10/019,114B

Input Set : A:\01997.013000 sequence listing.txt

Output Set: N:\CRF4\07302004\J019114B.raw

L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:1859 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1